

The Marginal Distribution of Compound Poisson INAR(1) Processes



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(Poisson) INAR(1) Processes

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Definition & Properties

Basic approach for **real-valued** processes:
stationary **ARMA(p,q) model**.

Let innovations $(\epsilon_t)_{\mathbb{Z}}$ be white noise, then

$$X_t = \alpha_1 \cdot X_{t-1} + \dots + \alpha_p \cdot X_{t-p} + \epsilon_t + \dots + \beta_q \cdot \epsilon_{t-q},$$

where $\alpha_1, \dots, \alpha_p, \beta_1, \dots, \beta_q \in \mathbb{R}$ suitably chosen.

Enumerous extension („ARMA alphabet soup“, Holan et al. (2010))
→ SARIMA, ARFIMA, VARMA, GARCH, ...

Not applicable to count data processes: generally, $\alpha \cdot X \notin \mathbb{N}_0$.

Several approaches of how to avoid “multiplication problem”.

Popular: Use appropriate thinning operations. (Weiß, 2008)

In this talk, we exclusively consider models based on

binomial thinning operator (Steutel & van Harn, 1979):

$$\alpha \circ X := \sum_{i=1}^X Y_i, \quad \text{where } Y_i \text{ are i.i.d. Bin}(1, \alpha),$$

i. e., $\alpha \circ X \sim \text{Bin}(X, \alpha)$ and has range $\{0, \dots, X\}$.

In particular, $E[\alpha \circ X] = E[\alpha \cdot X]$.

(\approx number of “survivors” from population of size X)

Let $(\epsilon_t)_{\mathbb{Z}}$ be i.i.d. with range $\mathbb{N}_0 = \{0, 1, \dots\}$,
denote $E[\epsilon_t] = \mu_\epsilon$, $V[\epsilon_t] = \sigma_\epsilon^2$. Let $\alpha \in (0; 1)$.

$(X_t)_{\mathbb{Z}}$ referred to as **INAR(1) process** if

$$\underbrace{X_t}_{\text{Population at time } t} = \underbrace{\alpha \circ X_{t-1}}_{\text{Survivors of time } t-1} + \underbrace{\epsilon_t}_{\text{Immigration}} .$$

plus approp. independence assumptions. (McKenzie, 1985)

Properties:

Homogeneous Markov chain with

$$\mathbb{P}(X_t = k \mid X_{t-1} = l) = \sum_{j=0}^{\min\{k,l\}} \binom{l}{j} \alpha^j (1 - \alpha)^{l-j} \cdot \mathbb{P}(\epsilon_t = k - j).$$

If INAR(1) process stationary (see below), then

$$\text{pgf}_X(z) = \text{pgf}_X(1 - \alpha + \alpha z) \cdot \text{pgf}_\epsilon(z).$$

In particular, if $\mu_\epsilon, \sigma_\epsilon < \infty$, we have

$$\mu_X = \frac{\mu_\epsilon}{1 - \alpha}, \quad \sigma_X^2 = \mu_X \cdot \frac{\frac{\sigma_\epsilon^2}{\mu_\epsilon} + \alpha}{1 + \alpha}.$$

(Note: X_t equidispersed iff ϵ_t equidispersed.)

Autocorrelation function: $\rho_X(k) = \alpha^k$, i. e., AR(1)-type.

For further properties and references, see Weiß (2008).

Most popular instance of INAR(1) family:

Poisson INAR(1) model,
$$X_t = \alpha \circ X_{t-1} + \epsilon_t.$$

Here, innovations $(\epsilon_t)_{\mathbb{Z}}$ i.i.d. $\text{Poi}(\lambda)$, such that $\mu_\epsilon = \sigma_\epsilon^2 = \lambda$.

Stationary marginal distribution:

also Poisson distribution, $\text{Poi}(\frac{\lambda}{1-\alpha})$, because:

additivity of Poisson distribution, and

Poisson distribution **invariant w.r.t. binomial thinning**:

If $X \sim \text{Poi}(\mu)$, then $\alpha \circ X \sim \text{Poi}(\alpha \cdot \mu)$.

Poisson INAR(1) has **equidispersed** marginals.

Modifications to get **overdispersed** marginals:

- modify thinning operation (Weiß, 2008), or
- modify distribution of innovations, e. g.,
negative binomially (NB) distributed ϵ_t 's.

In the following, we more generally consider
compound Poisson (CP) distributed innovations.



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Compound Poisson INAR(1) Processes

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Definition & Properties

Y_1, Y_2, \dots i.i.d. with range $\mathbb{N} = \{1, 2, \dots\}$,

denote pgf as $H(z)$ (**compounding distribution**).

Let $N \sim \text{Poi}(\lambda)$, independent of Y_1, Y_2, \dots

$\epsilon := Y_1 + \dots + Y_N$ **compound Poisson distributed:**

$\epsilon \sim \text{CP}(\lambda, H)$.

Then $\text{pgf}_\epsilon(z) = \exp(\lambda(H(z) - 1))$.

More precisely:

$\epsilon \sim \text{CP}_\nu(\lambda, H)$ if $H(z) = h_1 z + \dots + h_\nu z^\nu$ with $h_\nu > 0$,

$\epsilon \sim \text{CP}_\infty(\lambda, H)$ if Y_i have infinite range.

Properties:

- ϵ equidispersed iff $\nu = 1$: $CP_1(\lambda, H) = \text{Poi}(\lambda)$.
 ϵ overdispersed iff $\nu > 1$, e. g., ...
- $CP_\nu(\lambda, H)$ with $\nu < \infty$: **Hermite distribution** of order ν .
- $CP_\nu(\lambda, H)$ with $\nu < \infty$ and $h_x = 1/\nu$ for $x = 1, \dots, \nu$:
Poisson distribution of order ν , abbr. $\text{Poi}_\nu(\lambda)$.
- NB(n, π)-distribution: $CP_\infty(\lambda, H)$ with
 $\lambda := -n \ln \pi$ and $H(z)$ being pgf of log-series distribution:

$$H(z) = \frac{\ln(1 - (1 - \pi)z)}{\ln \pi} = \sum_{k=1}^{\infty} \frac{(1 - \pi)^k}{-k \ln \pi} z^k.$$

Properties: (cont.)

- **Additivity:**

If X_1, X_2 independent with $X_i \sim \text{CP}_\nu(\lambda_i, H_i)$ (common ν), then $X_1 + X_2$ $\text{CP}_\nu(\lambda, H)$ -distributed.

- **Invariance w.r.t. binomial thinning:**

If $X \sim \text{CP}_\nu(\lambda, H)$, then $\alpha \circ X \sim \text{CP}_\nu(\eta, G)$.

- Count data model parametrized by ν first factorial cumulants closed under addition and under binomial thinning *iff* it has CP_ν -distribution.

(Puig & Valero, 2007; Schweer & Weiß, 2014)

INAR(1) process $(X_t)_{t \in \mathbb{Z}}$ referred to as

CPINAR(1) process

if innovations $(\epsilon_t)_{\mathbb{Z}}$ i.i.d. $CP_\nu(\lambda, H)$ (possibly $\nu = \infty$).

$\nu = 1$: Poisson INAR(1) model.

Popular for $\nu = \infty$: NB-innovations, NB-INAR(1) model.

Heathcote (1966), Schweer & Weiß (2014):

If $H'(1) < \infty$, then unique **stationary** marginal distribution of $(X_t)_{t \in \mathbb{Z}}$ is $CP_\nu(\eta, G)$, where

$$\eta(G(z) - 1) = \lambda \sum_{i=0}^{\infty} (H(1 - \alpha^i + \alpha^i z) - 1).$$



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Marginal Distribution

See above:

If INAR(1)'s innovations $(\epsilon_t)_{\mathbb{Z}}$ CP_{ν} -distributed (possibly $\nu = \infty$), then INAR(1)'s observations $(X_t)_{t \in \mathbb{Z}}$ also CP_{ν} -distributed, with same compounding order ν .

E.g., for the purpose of likelihood computations:

How to compute **efficiently and precisely**
pmf of **marginal distribution** of $(X_t)_{t \in \mathbb{Z}}$?

If compounding order **finite**, $\nu < \infty$,

then marginal distribution can be computed exactly!

Scheme: (see Weiß & Puig (2015) for details & references)

1. Compute parameters η and g_1, \dots, g_ν of marginal distribution:

(a) If innovations' param. λ and h_1, \dots, h_ν readily available, solve following linear equations in η and g_1, \dots, g_ν :

$$g_1 + \dots + g_\nu = 1, \quad \frac{\lambda}{\eta} - (1 - \alpha) \cdot g_1 - \dots - (1 - \alpha)^\nu \cdot g_\nu = 0,$$

$$h_k \cdot \frac{\lambda}{\eta} - (1 - \alpha^k) \cdot g_k + \alpha^k \sum_{i=k+1}^{\nu} \binom{i}{k} (1 - \alpha)^{i-k} \cdot g_i = 0, \quad k = \nu, \dots, 2.$$

(b) If innovations' CP_ν -distr. via fact. cumul. $\kappa_{(1), \epsilon}, \dots, \kappa_{(\nu), \epsilon}$,

then first $\kappa_{(1), X}, \dots, \kappa_{(\nu), X}$ via $\kappa_{(n), X} = \frac{\kappa_{(n), \epsilon}}{1 - \alpha^n}$,

afterwards expand $\eta(G(z) - 1) := \sum_{r=1}^{\nu} \frac{\kappa_{(r), X}}{r!} \cdot (z - 1)^r$.

Scheme: (see Weiß & Puig (2015) for details & references)

1. (...)

2. Recursive scheme for computation of pmf $P(X = k)$:

$$P(X = 0) = e^{-\eta}, \quad (\text{“Panjer recursion”})$$

$$P(X = k) = \frac{\eta}{k} \cdot \sum_{j=1}^{\min\{k, \nu\}} j g_j \cdot P(X = k - j) \quad \text{for } k \geq 1.$$

In the case $\nu = \infty$, however, Step 1 is crucial point.

Idea: Replace infinite compounding structure by finite one, then continue with above Scheme.

Hermite- ν approximation, type 1:

For innovations ϵ_t being $CP_\infty(\lambda, H)$ -distributed,
define ν -th order approximation $CP_\nu(\tilde{\lambda}, \tilde{H})$ by

$$\tilde{\lambda} := \lambda, \quad \tilde{h}_1 := h_1, \quad \dots, \quad \tilde{h}_{\nu-1} := h_{\nu-1}, \quad \tilde{h}_\nu := 1 - \sum_{k=1}^{\nu-1} h_k.$$

Then proceed according to Steps 1(a) and 2 of above Scheme.

Preserves $P(\epsilon = 0), \dots, P(\epsilon = \nu - 1)$,

approximates remaining innovations' probabilities.

1st order approximation = Poisson approximation.

Hermite- ν approximation, type 2:

For innovations ϵ_t being $CP_\infty(\lambda, H)$ -distributed with mean μ_ϵ , define ν -th order approximation $CP_\nu(\tilde{\lambda}, \tilde{H})$ by

$$\tilde{h}_1 := \frac{h_1}{\sum_{k=1}^{\nu} h_k}, \dots, \tilde{h}_\nu := \frac{h_\nu}{\sum_{k=1}^{\nu} h_k}, \quad \tilde{\lambda} := \frac{\mu_\epsilon}{\sum_{k=1}^{\nu} k \tilde{h}_k}.$$

Then proceed according to Steps 1(a) and 2 of above Scheme.

Preserves observations' mean μ_X ,

approximates all innovations' probabilities.

1st order approximation = Poisson approximation.

Performance evaluation:

NB-INAR(1) model with $NB(n, \pi)$ -distributed innovations,
different levels of mean parameter n ,
dispersion parameter π , autocorrelation parameter α .

Benchmark:

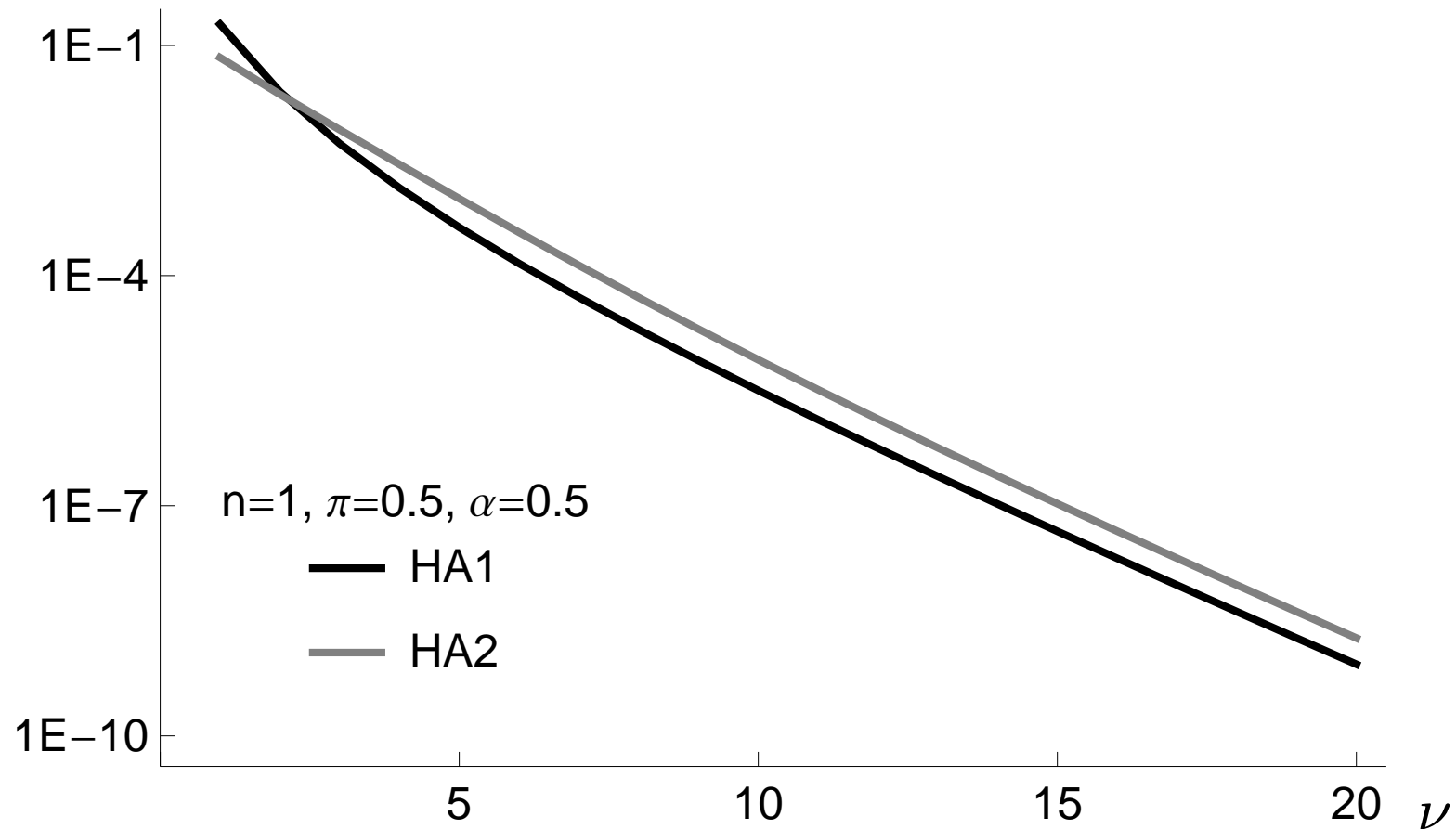
Numerically exact Markov chain approach (Weiß, 2010).

Concerning computational **efficiency**:

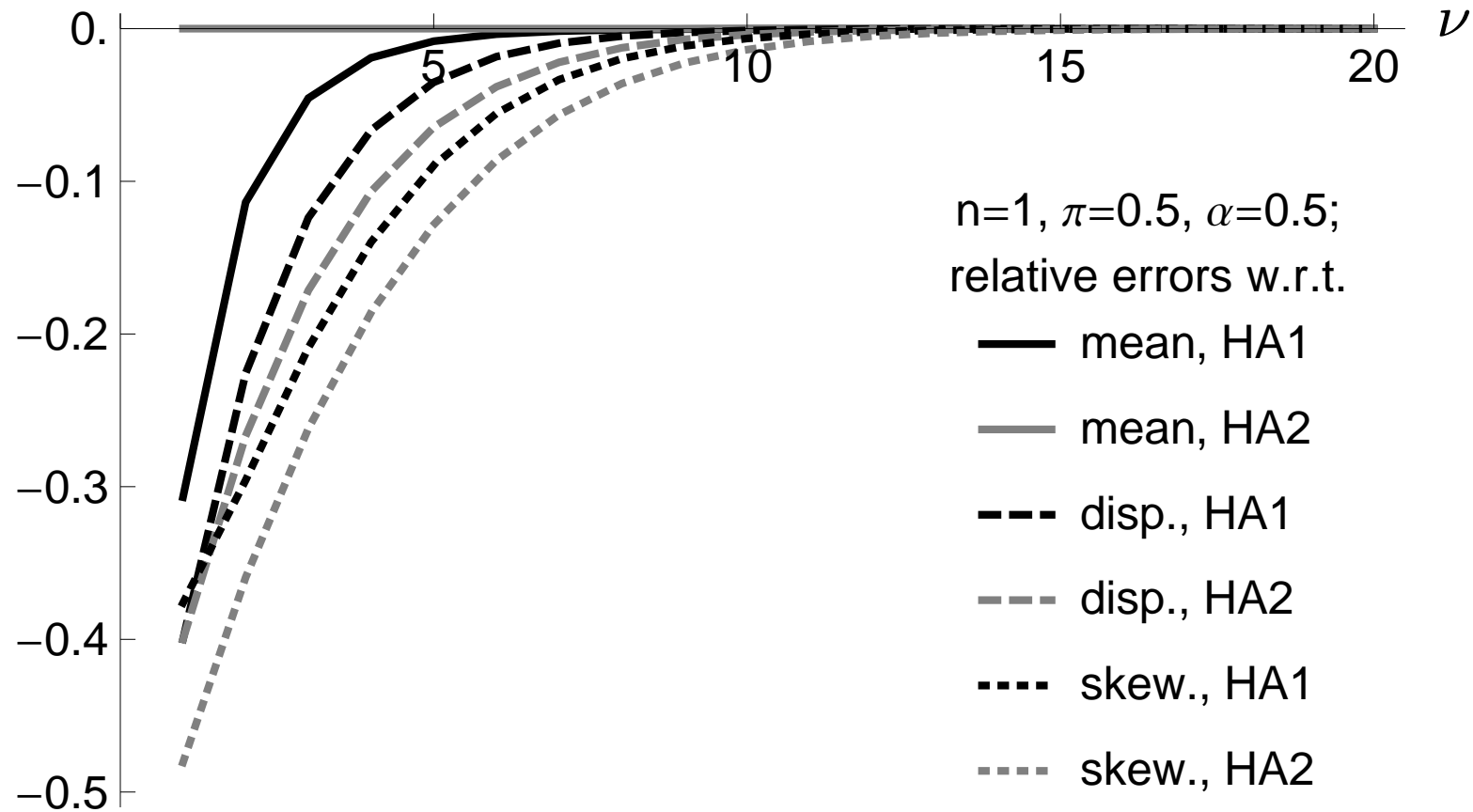
Computing time for approximations: < 0.1 s.

Computing time for MC approach: 70 – 100 s.

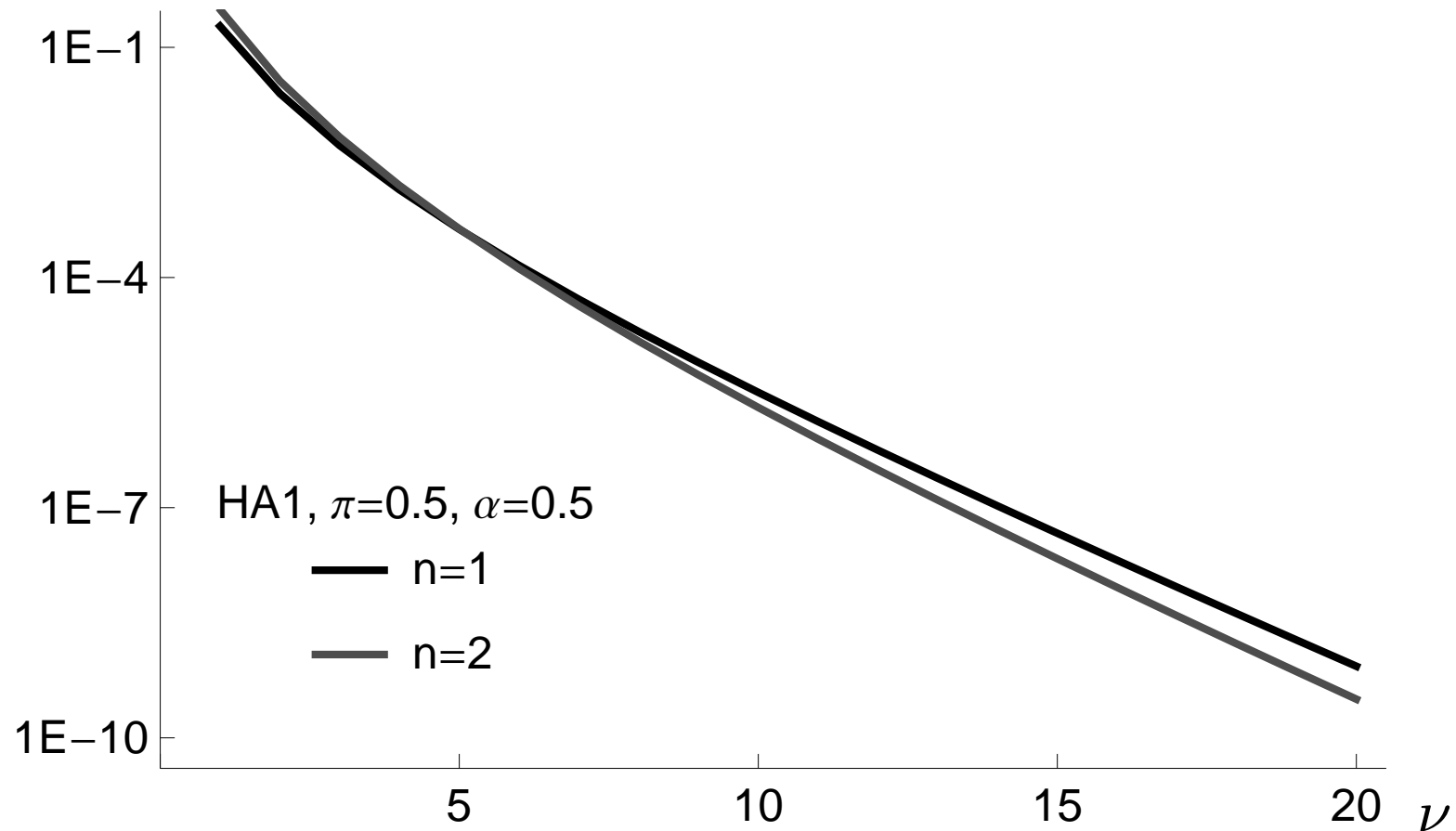
Overall quality of approximation via Kullback-Leibler divergence:



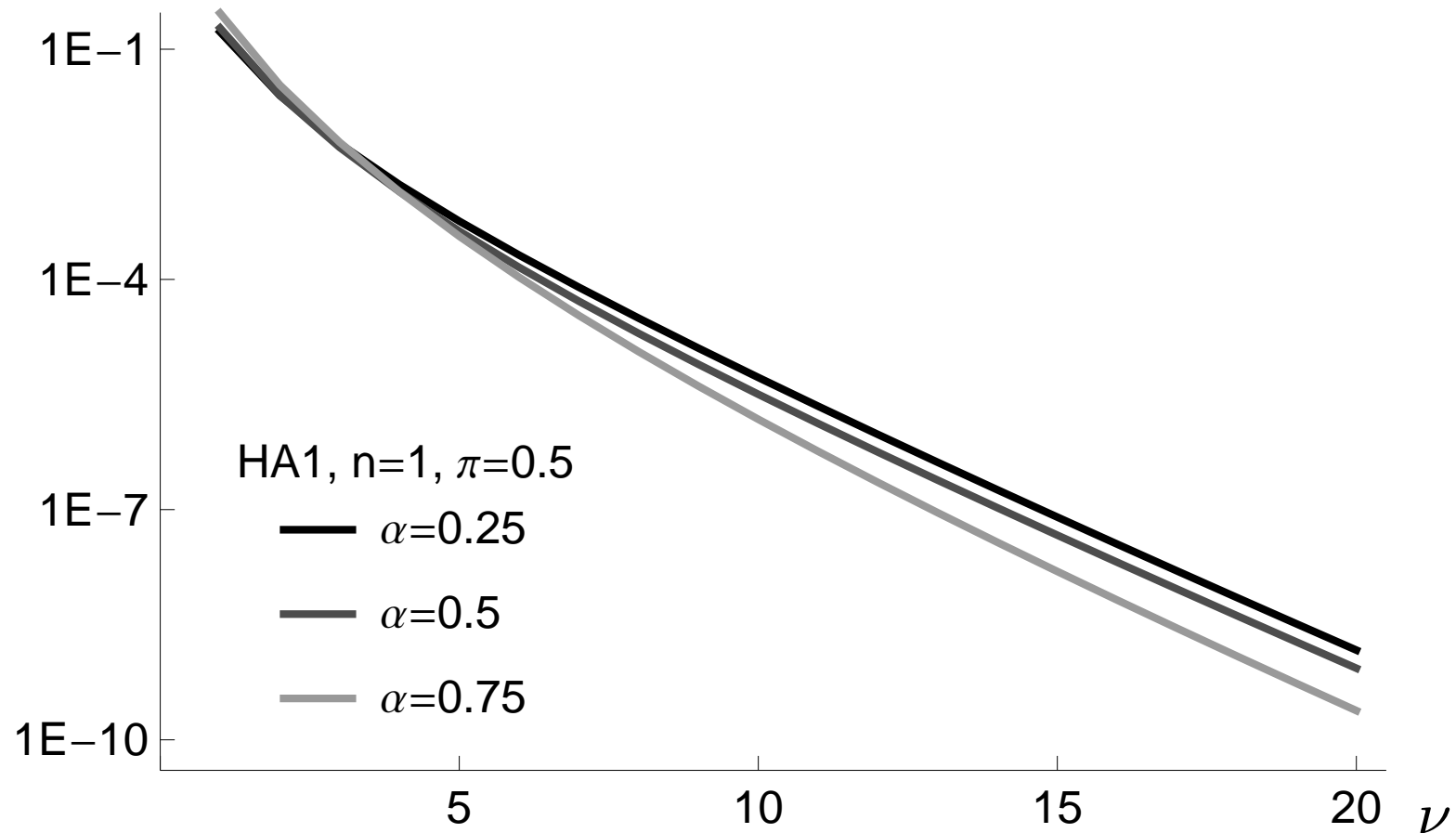
Relative errors $\frac{\text{approx} - \text{true}}{\text{true}}$ for mean, dispersion index and skewness:



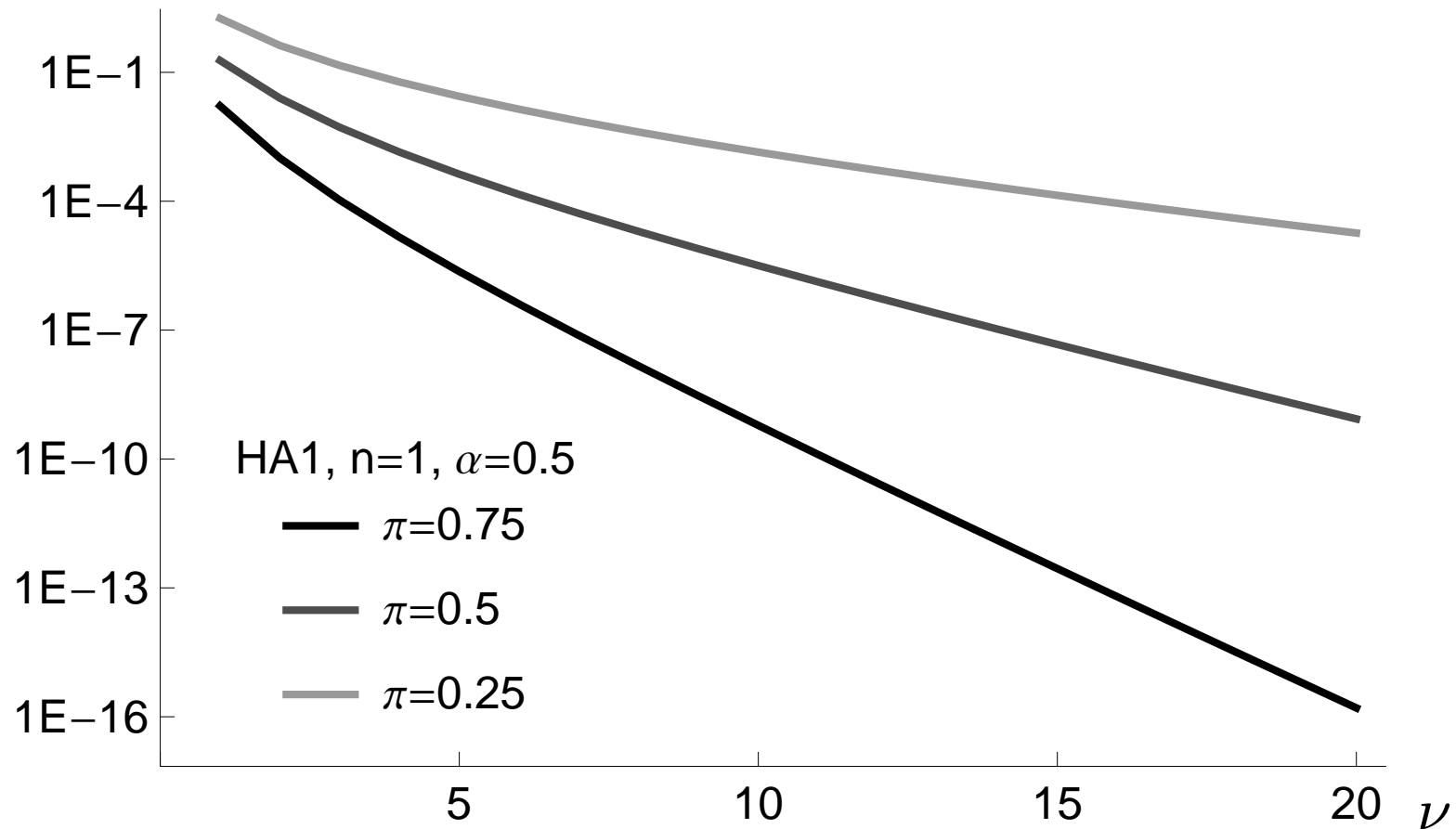
Overall quality of type 1 approximation via KL divergence:



Overall quality of type 1 approximation via KL divergence:



Overall quality of type 1 approximation via KL divergence:





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Conclusions

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... and Future Research

- Type 1 approximation performs best concerning KL divergence, and concerning dispersion index or skewness.
- If one chooses $\nu \geq 15$, then difference between exact distribution and type 1 approximation negligible.
- Both approximations much faster than MC approach.

Future research:

empirical version of the Hermite approximation, e. g.,
based on first ν empirical factorial cumulants.

(\rightarrow semi-parametric estimation of INAR(1)'s marginal)

Thank You for Your Interest!



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